

200

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83593563.q MEAAELVKKIPPELEEGCAPDKRKEISRIVPPFRGARRPTILPAQ-QRR-TALPQAAAAPSRLQLQVRGA
74530011.q MEEAELVKKIPPELEEGCAPDKRKEISRIVPPFRGARRPTILPAQ-QRR-TALPQAAAAPSRLQLQVRGA
22327668.p MENPNPPAVKLKKKIPELEESQEDIKRKEMSKIKVSAEIKRSHSSSPKRSPRRNSGEGTPILWRKTGAASFRHASP-LRKESHSHDGVAAGGGDGPSAKPTDKQYINILQSMQAQAVHVDLNGQOIIINSMAEAKLYCFSAAEAKGKDSINILWDGDAAVAKNIFCRCSSESTGCEFVKRN
18408889.p MVKLLQDPPITNKKLKKMPELEKSKQEDIKRKEMSRICKVSIEIKRKE
15230755.p MEEELKKKMPPELEOSQEDIKRKEMSRICKLSNEIRCPSPVILPRRPLR-RIQCSMNSN-PSECKPTDKQYINILQSMQAQAVHVDLNGQOIIINSMAEAKLYCFSAAEAVCQNEIDVMVDRAPAMMIAACLSNGESTGKFVKRN
15230754.p MENP-PAEELKKKIDPELEESQEDIKRKEMSRICKVSTEIRCRSHSVPHPARRNIGEGAPSWRKSGAASFRNAPLKRERIQNSMRRLSEVGGGGSPAKPTDKQYINILQSMQAQAVHADLNGQOIIINSMAEAKVYCYSAEAVCQNEIDVMVDRAPAMMIAACLSNGESTGKFVKRN
15230753.p lias_TGFBR

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83593563.q SCKEPNIVASPTPHDDC-LVYLICLSTDTR-TPEETISGPESICARPHPESS-KPRLVOLNGS-KSGCLNNVCSDPSPLOSTIASLTNLATVKINKVRSRVRAENIER-ECCGSEPSD-DAKEEPSSCTTREGAPR-GATAPESSSGKAKNSDS-ECKVGFRRILSSKAPAL
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22327668.p MGERFSWVTTISCFYDDDCGILIIITNDISALFQCRPPAKRNWQE-CDSDFCRGTINGVASRLGEDESEAUVSKLGDSQPIOAATASKISIDLASKVINKVRSKMRADNAASHP-ECCGSEPSD-SDQGFEADAFTSDREDAETINDASTRGNDIQCSPGVPLCNDDKSSKASCESENDDRNVVVEKKLTSKAEE
18408889.p TGEXFSWVTTISCFYDDDCGILIIITSDIAVAPYINPRLSLERLKPOE-PK-CLGDSKCAVISKGHDSPQIOVATAKISIDLASKVINKVRSKMRADNAASHP-ECCGSEPSD-SDQGFEADAFTSDREDAETINDASTRGNDIQCSPGVPLCNDDKSSKASCESENDDRNVVVEKKLTSKAEE
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15230753.p lias_TGFBR

600

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83593563.q WANRGIPWPRGHEND-ACKNR-TNLPHQEFHQENGQSHKEVPEPINDPQDQIEFVQEVKVEVSCSW-SEN-ASTSSMSTGSTNSAAERIDREADCLDFEIL-EDIAIKEQVQGSCCTVYHGL-YGSDVAVKVFSKQYESEEVIL-TFROEVSLMKLRHPNILLFCMCAVTSPQ-RLCIVTEFPLRGSLFRL
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lias_TGFBR

800

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83593563.q LOKNAGTKLDRRRVHMAIDTACRMNYLHN-SSPPIVHRDLKSSNLLVDKN-TVKVADFGCLSR-LKGETFLTKTKGKTPQ-MAEVLVNE-SNPE-NYGVIL-BLTOKEIP-BNLNTM0V1GAVGFMNMRLEIP-SEDPQNTSLILSCWE
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lias_TGFBR

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83593563.q VFNKPILSASSDQ-RCQELDQDQYQVQNLQALQ-QLR-TAAAKMSVDC-
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18408889.p DAKLPRTFQELMERLDRDQYKMICFOATR-ALSDKNCEREPKERVGLFEOQQQLVETCCAGVPMSSSTIEQGYGIEINIMKVG-
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